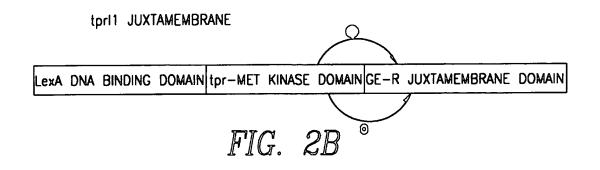
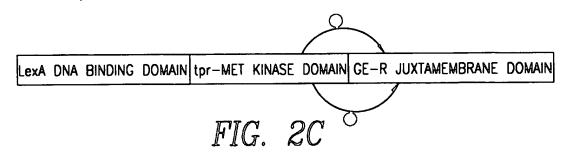


FIG. 2A



tpri1 C-TERMINAL



i

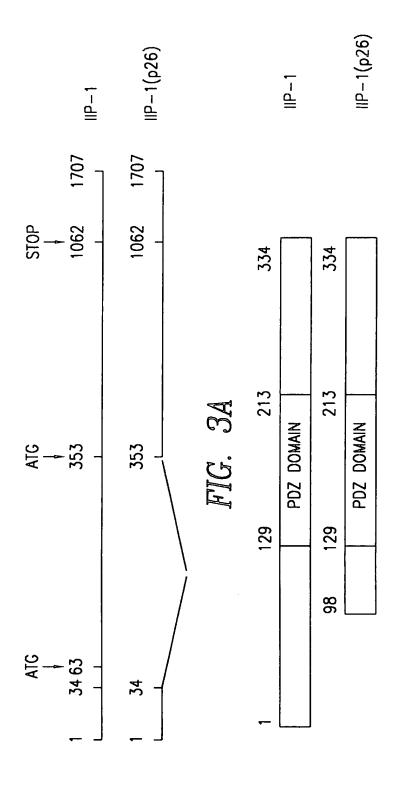
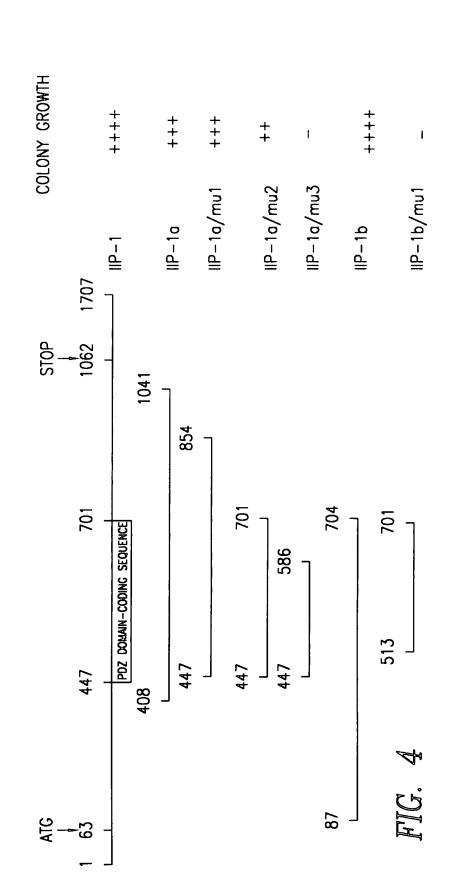


FIG. 3B



## Fig. 5

- 1 MSRPRRLAG TSGSDKGLSG KRTKTENSGE ALAKVEDSNP QKTSATKNCL
- KNLSSHWLMK SEPESRLEKG VDVKFSTEDL KAQPKQTTCW DGVRNYQARN 51
- + FLRAMKLGEE AFFYHSNCKE PGIAGLMKIV KEAYPDHTÖF EKNNPHYDPS 101
- \*KEDNPKWSM VDVQFVRMMK RFIPLAELKS YHQAHKATGG PLKNMVLFTR 151
- 201 QRLSIQPLTQ EEFDFVLSLE EKEPS\*
- ▼ putative N-glycosylation site
- putative Casein kinase II phosphorylation site **\Q**
- putative N-myristoylation site
- putative Protein kinase C phosphorylation site 
   ✓ putative Tyrosine kinase phosphorylation site
   ✓ putative Protein kinase C phosphorylation site
- **←→** putative NLS

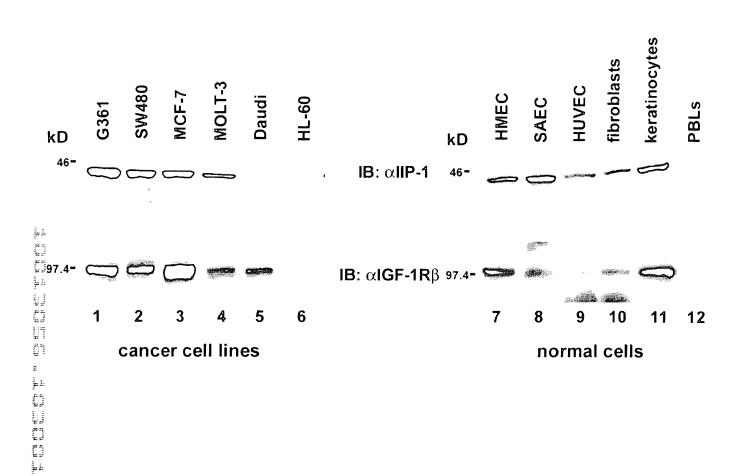






Fig. 7

